

1638

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#18

## RAW SEQUENCE LISTING

DATE: 11/13/2002

PATENT APPLICATION: US/09/251,638B

TIME: 13:49:23

Input Set : A:\EP.txt

Output Set: N:\CRF4\11132002\I251638B.raw

3 <110> APPLICANT: DANIELL, HENRY  
 5 <120> TITLE OF INVENTION: GENETIC ENGINEERING OF COTTON TO INCREASE FIBER  
 6 STRENGTH, WATER ABSORPTION AND DYE BINDING  
 8 <130> FILE REFERENCE: 1483-R-00  
 10 <140> CURRENT APPLICATION NUMBER: 09/251,638B  
 11 <141> CURRENT FILING DATE: 1999-02-17  
 13 <150> PRIOR APPLICATION NUMBER: 60/074,997  
 14 <151> PRIOR FILING DATE: 1998-02-17  
 16 <160> NUMBER OF SEQ ID NOS: 5  
 18 <170> SOFTWARE: PatentIn Ver. 2.1  
 20 <210> SEQ ID NO: 1  
 21 <211> LENGTH: 5  
 22 <212> TYPE: PRT  
 23 <213> ORGANISM: Artificial Sequence  
 25 <220> FEATURE:  
 26 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic  
 27 peptide  
 29 <400> SEQUENCE: 1  
 30 Val Pro Gly Val Gly  
 31 1 5  
 34 <210> SEQ ID NO: 2  
 35 <211> LENGTH: 5  
 36 <212> TYPE: PRT  
 37 <213> ORGANISM: Artificial Sequence  
 39 <220> FEATURE:  
 40 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic  
 41 peptide  
 43 <400> SEQUENCE: 2  
 44 Gly Val Gly Val Pro  
 45 1 5  
 48 <210> SEQ ID NO: 3  
 49 <211> LENGTH: 605  
 50 <212> TYPE: PRT  
 51 <213> ORGANISM: Artificial Sequence  
 53 <220> FEATURE:  
 54 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic  
 55 peptide  
 57 <400> SEQUENCE: 3  
 58 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly  
 59 1 5 10 15  
 61 Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val  
 62 20 25 30  
 64 Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly

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65          35          40          45
67 Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val
68          50          55          60
70 Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro
71 65          70          75          80
73 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly
74          85          90          95
76 Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val
77          100         105         110
79 Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly
80          115         120         125
82 Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val
83          130         135         140
85 Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro
86 145         150         155         160
88 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly
89          165         170         175
91 Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val
92          180         185         190
94 Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly
95          195         200         205
97 Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val
98          210         215         220
100 Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro
101 225         230         235         240
103 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly
104          245         250         255
106 Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val
107          260         265         270
109 Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly
110          275         280         285
112 Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val
113          290         295         300
115 Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro
116 305         310         315         320
118 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly
119          325         330         335
121 Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val
122          340         345         350
124 Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly
125          355         360         365
127 Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val
128          370         375         380
130 Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro
131 385         390         395         400
133 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly
134          405         410         415
136 Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val
137          420         425         430

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139 Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly
140           435                     440                     445
142 Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val
143           450                     455                     460
145 Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro
146 465                     470                     475                     480
148 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly
149           485                     490                     495
151 Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val
152           500                     505                     510
154 Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly
155           515                     520                     525
157 Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val
158           530                     535                     540
160 Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro
161 545                     550                     555                     560
163 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly
164           565                     570                     575
166 Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val
167           580                     585                     590
169 Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro
170           595                     600                     605
173 <210> SEQ ID NO: 4
174 <211> LENGTH: 100
175 <212> TYPE: PRT
176 <213> ORGANISM: Artificial Sequence
178 <220> FEATURE:
179 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
180 peptide
182 <400> SEQUENCE: 4
183 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly
184   1           5           10           15
186 Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val
187           20           25           30
189 Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly
190           35           40           45
192 Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val
193           50           55           60
195 Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro
196  65           70           75           80
198 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly
199           85           90           95
201 Val Gly Val Pro
202           100
206 <210> SEQ ID NO: 5
207 <211> LENGTH: 605
208 <212> TYPE: PRT
209 <213> ORGANISM: Artificial Sequence
211 <220> FEATURE:

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212 <221> NAME/KEY: repeat_unit
213 <222> LOCATION: 1..605
214 <223> OTHER INFORMATION: Repeats at least once
216 <220> FEATURE:
217 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
218     peptide
220 <400> SEQUENCE: 5
221 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly
222   1           5           10           15
224 Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val
225           20           25           30
227 Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly
228           35           40           45
230 Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val
231           50           55           60
233 Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro
234   65           70           75           80
236 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly
237           85           90           95
239 Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val
240           100          105          110
242 Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly
243           115          120          125
245 Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val
246           130          135          140
248 Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro
249   145          150          155          160
251 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly
252           165          170          175
254 Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val
255           180          185          190
257 Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly
258           195          200          205
260 Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val
261           210          215          220
263 Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro
264   225          230          235          240
266 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly
267           245          250          255
269 Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val
270           260          265          270
272 Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly
273           275          280          285
275 Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val
276           290          295          300
278 Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro
279   305          310          315          320
281 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly
282           325          330          335

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```

284 Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val
285           340           345           350
287 Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly
288           355           360           365
290 Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val
291       370           375           380
293 Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro
294 385           390           395           400
296 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly
297           405           410           415
299 Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val
300           420           425           430
302 Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly
303           435           440           445
305 Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val
306       450           455           460
308 Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro
309 465           470           475           480
311 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly
312           485           490           495
314 Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val
315           500           505           510
317 Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly
318           515           520           525
320 Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val
321       530           535           540
323 Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro
324 545           550           555           560
326 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly
327           565           570           575
329 Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val
330           580           585           590
332 Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro
333           595           600           605

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**VERIFICATION SUMMARY**

DATE: 11/13/2002

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TIME: 13:49:24

Input Set : A:\EP.txt

Output Set: N:\CRF4\11132002\I251638B.raw